

LISTING OF CLAIMS:

1. - 132. (Canceled)

133. (Currently amended) A method for identifying a chromosomal deletion,
comprising the steps of :

~~DNA probe set, said probe set comprising a first probe set and a second probe set,~~
~~hybridizing a first probe set to a chromosome, said first probe set being sufficient in~~
~~length and substantially complementary to a deletion region of a chromosome but~~
~~less than an entire chromosome such that said first probe set will hybridize to the~~
~~deletion region if it is present, and~~

hybridizing a second probe set to a chromosome, said second probe set being
sufficient in length and substantially complementary to nucleotides in the regions of
the chromosome flanking the deletion region but less than an entire chromosome
such that said second probe set will hybridize to the flanking regions regardless of
whether the deletion region is present;

wherein said first and second probe sets are detectably labeled, and wherein the
label for the first probe set is distinguishable from the label for the second probe set.

134 - 143. (Canceled)

144. (New) The method of claim 133, further comprising adding blocking nucleic acid that comprises fragments which are substantially complementary to repetitive segments in the targeted chromosomal DNA.